

501 TEC 475860

SEQUENCE LISTING

<110> Allen, Steve  
Lee, Jian Ming

<120> Plant Protein Kinases

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<150> 60/092, 438

<151> July 10, 1998

<160> 23

<170> Microsoft Office 97

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<213> Zea mays

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TOMATO - PEACH SAB

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**3/3**

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35 40 45

Ala Thr Gly Lys Glu Phe Ala Cys Lys Ser Ile Leu Lys Xaa Leu Val  
50 55 60

Thr Asp Asp Asp Val Glu Asp Val Arg Arg Glu Ile Gln Ile Met His  
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Asp Ala Val Ala Val  
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<213> Oryza sativa

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 35 40 45

Val Gly Thr Pro Arg Arg Arg Gly Ser Lys Ser Gly Ser Thr Thr Pro  
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Gly His Gln Thr Pro Gly Val Ala Trp Pro Ser Pro Tyr Pro Ser Gly  
 65 70 75 80

Gly Ala Ser Pro Leu Pro Ala Gly Val Ser Pro Ser Pro Ala Arg Ser  
 85 90 95

Thr Pro Arg Arg Phe Phe Lys Arg Pro Phe Pro Pro Pro Ser Pro Ala  
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Lys His Ile Lys Ala Thr Leu Ala Lys Arg Leu Gly Gly Lys Pro  
 115 120 125

Lys Glu Gly Thr Ile Pro Glu Glu Gly Val Gly Ala Gly Gly Gly  
 130 135 140

Gly Gly Gly Ala Ala Asp Gly Ala Glu Thr Glu Arg Pro Leu Asp Lys  
 145 150 155 160

Thr Phe Gly Phe Ser Lys Asn Phe Gly Ala Lys Tyr Glu Leu Gly Lys  
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Glu Val Gly Arg Gly His Phe Gly His Thr Cys Ser Ala Val Val Lys  
180 185 190

Lys Gly Glu Tyr Lys Gly Gln Thr Val Ala Val Lys Ile Ile Ala Lys  
195 200 205

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Lys Ile Leu Arg Ala Leu Ser Gly His Asn Asn Leu Val Lys Phe Tyr  
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Asp Ala Cys Glu Asp Gly Leu Asn Val Tyr Ile Val Met Glu Leu Cys  
245 250 255

Glu Gly Gly Glu Leu Leu Asp Arg Ile Leu Ala Arg Gly Gly Arg Tyr  
260 265 270

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275 280 285

Ala Phe Cys His Leu Gln Gly Val Val His Arg Asp Leu Lys Pro Glu  
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Asp Phe Gly Leu Ser Asp Phe Ile Arg Pro Asp Glu Arg Leu Asn Asp  
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Ile Val Gly Ser Ala Tyr Tyr Val Ala Pro Glu Val Leu His Arg Ser  
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Tyr Ser Met Glu Ala Asp Ile Trp Ser Ile Gly Val Ile Thr Tyr Ile  
355 360 365

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370 375 380

Phe Arg Ser Val Leu Arg Ala Asp Pro Asn Phe Asp Asp Ser Pro Trp  
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Pro Thr Val Ser Ala Glu Ala Lys Asp Phe Val Lys Arg Phe Leu Asn  
405 410 415

Lys Asp Tyr Arg Lys Arg Met Thr Ala Val Gln Ala Leu Thr His Pro  
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Trp Leu Arg Asp Glu Gln Arg Gln Ile Pro Leu Asp Ile Leu Ile Phe  
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Leu Lys Ala Leu Ser Lys Ala Leu Arg Glu Asp Glu Leu Leu Tyr Leu  
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Lys Leu Gln Phe Lys Leu Leu Glu Pro Arg Asp Gly Phe Val Ser Leu  
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Asp Asn Phe Arg Thr Ala Leu Thr Arg Tyr Leu Thr Asp Ala Met Lys  
500 505 510

Glu Ser Arg Val Leu Glu Phe Leu His Ala Leu Glu Pro Leu Ala Tyr  
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Arg Arg Met Asp Phe Glu Glu Phe Cys Ala Ala Ala Ile Ser Pro Tyr  
530 535 540

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Gln Gln Phe Glu Gln Glu Gly Asn Arg Val Ile Ser Val Glu Glu Leu  
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Ala Gln Glu Leu Asn Leu Ala Pro Thr His Tyr Ser Ile Val Gln Asp  
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20 25 30

POINT TO PROTEIN

Pro Pro Ser Pro Ala Lys His Ile Arg Ala Leu Leu Ala Arg Xaa His  
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Gly Ser Val Lys Pro Asn Glu Ala Ser Ile Pro Glu Ala Ser Xaa Cys  
50 55 60

Glu Leu Gly Leu Asp Lys Ser Phe Gly Phe Ala Lys Gln Phe Ser Ala  
65 70 75 80

His Tyr Glu Leu Ser Asp Glu Xaa Gly Arg Gly His Phe Gly Tyr Thr  
85 90 95

Cys Ser Ala Lys Gly Lys Gly Ala Phe Lys Gly Leu Asn Val Ala  
100 105 110

Val Lys Val Ile Pro Lys Ala Lys Met Thr Thr Ala Ile Ala Ile Glu  
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Phe Ser Ser Lys Glu Glu Asn Ser Pro Leu Lys Val Ile Asp Phe Gly  
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Leu Ser Asp Phe Val Lys Pro Asp Glu Arg Leu Asn Asp Ile Val Gly  
65 70 75 80

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<212> DNA

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35 40 45  
Asn Gly Ala Glu Pro Gly His Ile Ile Val Thr Ser Ile Asp Gly Arg  
50 55 60  
Asn Gly Gln Ala Lys Gln Thr Ile Ser Tyr Met Ala Glu Arg Val Val  
65 70 75 80  
Gly His Gly Ser Phe Gly Thr Val Phe Gln Ala Lys Cys Leu Glu Thr  
85 90 95  
Gly Glu Thr Val Ala Ile Lys Lys Val Leu Gln Asp Lys Arg Tyr Lys  
100 105 110  
Asn Arg Glu Leu Gln Thr Met Arg Val Leu Asp His Pro Asn Val Val  
115 120 125  
Ala Leu Lys His Cys Phe Phe Ser Lys Thr Glu Lys Glu Glu Leu Tyr  
130 135 140  
Leu Asn Leu Val Leu Glu Tyr Val Pro Glu Thr Ala His Arg Val Ile  
145 150 155 160  
Lys His Tyr Asn Lys Met Asn Gln Arg Met Pro Leu Ile Tyr Ala Lys  
165 170 175  
Leu Tyr Met Tyr Gln Ile Cys Arg Ala Leu Ala Tyr Ile His Asn Ser  
180 185 190  
Ile Gly Val Cys His Arg Asp Ile Lys Pro Gln Asn Leu Leu Val Asn  
195 200 205  
Pro His Thr His Gln Leu Lys Leu Cys Asp Phe Gly Ser Ala Lys Val  
210 215 220  
Leu Val Lys Gly Glu Pro Asn Ile Ser Tyr Ile Cys Ser Arg Tyr Tyr  
225 230 235 240  
Arg Ala Pro Glu Leu Ile Phe Gly Ala Thr Glu Tyr Thr Ala Ile  
245 250 255  
Asp Val Gly Ser Ala Gly Cys Val Leu Ala Glu Leu Leu Gly Gln  
260 265 270  
Pro Leu Phe Pro Gly Glu Ser Gly Val Asp Gln Leu Val Glu Ile Ile  
275 280 285

Lys Val Leu Gly Thr Pro Thr Arg Glu Glu Ile Lys Cys Met Asn Pro  
290 295 300

Asn Tyr Thr Glu Phe Lys Phe Pro Gln Ile Lys Ala His Pro Trp His  
305 310 315 320

Lys Ile Phe His Lys Arg Met Pro Ala Glu Ala Val Asp Leu Val Ser  
325 330 335

Arg Leu Leu Gln Tyr Ser Pro Lys Leu Arg Ser Thr Ala Leu Glu Ala  
340 345 350

Leu Val His Pro Phe Phe Asp Glu Leu Arg Asp Pro Asn Thr Arg Leu  
355 360 365

Pro Asn Gly Arg Phe Leu Pro Pro Leu Phe Asn Phe Lys Pro His Glu  
370 375 380

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caccacatg ggtgctgaga agttgcctga tcagatgcat gatctgaaga taaggacga 300  
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TOMATO PROTEIN SEQUENCES

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Thr Ser Met Gly Ala Glu Lys Leu Pro Asp Gln Met His Asp Leu Lys  
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Ile Arg Asp Asp Lys Glu Xaa Glu Xaa Xaa Thr Ile Ile Asn Xaa Xaa  
35 40 45

Gly Thr Glu Xaa Gly His Ile Ile Val Thr Thr Gly Gly Xaa Asn  
50 55 60

Gly Xaa Pro Lys Xaa Thr Val Ser Tyr Met Ala Xaa Arg Ile Val Gly  
65 70 75 80

Gln Gly Ser Phe Gly Ile Val Phe Gln Ala Lys Phe Trp Arg Gln Gly  
85 90 95

Glu Thr Val Ala Ile Lys Xaa Val Leu  
100 105

<210> 13  
<211> 1429  
<212> DNA  
<213> Glycine max

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<222> (1202)

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*Glycine max*  
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cggttgtgt ggcaccaact tcgggttga gagaaggccag tggcatgga gcagcaggtg 180  
ttgatagatt gccagaggag atgaacgata tgaaaattag ggtatgataga gaaatggaag 240  
ccacagttgt tgatggcaac ggaacggaga caggacatat cattgtgact accatgggg 300  
gtagaaatgg tcagcccaag cagactataa gctacatggc agagcgtgtt gtagggcatg 360  
gatcatttgg agttgtctc caggctaagt gcttggaaac cggtgaaact gtggctatca 420  
aaaaggttct tcaagacaag aggtacaaga accgggagct gcaaacaatg cgccttctg 480  
accacccaaa tgtcggtgct ttgaagcact gtttctttc aaccactgaa aaggatgaac 540  
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acaacaagtt taaccaaagg atgccaatgttatatgtgaa actctataca taccagatct 660  
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tgccgtttct gcttgtaan tgtacaac tgaagtgttgc ttcatataga atgcgngctt 1380  
cctcattaaa gaaattgtgg accttatgan tcgtnccgt aacagttag 1429

<210> 14  
<211> 399  
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<213> Glycine max

*Aut*  
*3/3*

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<222> (391)

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Met Thr Ser Val Gly Val Ala Pro Thr Ser Gly Leu Arg Glu Ala Ser  
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Gly His Gly Ala Ala Gly Val Asp Arg Leu Pro Glu Glu Met Asn Asp  
35 40 45

Met Lys Ile Arg Asp Asp Arg Glu Met Glu Ala Thr Val Val Asp Gly  
50 55 60

Asn Gly Thr Glu Thr Gly His Ile Ile Val Thr Thr Ile Gly Gly Arg  
65 70 75 80

Asn Gly Gln Pro Lys Gln Thr Ile Ser Tyr Met Ala Glu Arg Val Val  
85 90 95

Gly His Gly Ser Phe Gly Val Val Phe Gln Ala Lys Cys Leu Glu Thr  
100 105 110

Gly Glu Thr Val Ala Ile Lys Lys Val Leu Gln Asp Lys Arg Tyr Lys  
115 120 125

Asn Arg Glu Leu Gln Thr Met Arg Leu Leu Asp His Pro Asn Val Val  
130 135 140

Ala Leu Lys His Cys Phe Phe Ser Thr Thr Glu Lys Asp Glu Leu Tyr  
145 150 155 160

Leu Asn Leu Val Leu Glu Tyr Val Pro Glu Thr Val Asn Arg Val Ile  
165 170 175

Lys His Tyr Asn Lys Phe Asn Gln Arg Met Pro Leu Ile Tyr Val Lys  
180 185 190

Leu Tyr Thr Tyr Gln Ile Phe Arg Ala Leu Ser Tyr Ile His Arg Cys  
195 200 205

Ile Gly Val Cys His Arg Asp Ile Lys Pro Gln Asn Leu Leu Val Asn  
210 215 220

Pro His Thr His Gln Val Lys Leu Cys Asp Phe Gly Ser Ala Lys Val  
225 230 235 240

Leu Val Lys Gly Glu Pro Asn Ile Ser Tyr Ile Cys Ser Arg Tyr Tyr  
245 250 255

Arg Ala Pro Glu Leu Ile Phe Gly Ala Thr Glu Tyr Thr Thr Ala Ile  
260 265 270

Asp Val Trp Ser Val Gly Cys Val Leu Ala Glu Leu Leu Gly Gln  
275 280 285

Pro Leu Phe Pro Gly Glu Ser Gly Val Asp Gln Leu Val Glu Ile Ile  
290 295 300

Lys Val Leu Gly Thr Pro Thr Arg Glu Glu Ile Lys Cys Met Asn Pro  
305 310 315 320

Asn Tyr Thr Glu Phe Lys Phe Pro Gln Ile Lys Ala His Pro Trp His  
325 330 335

Lys Ile Phe His Lys Arg Met Pro Pro Glu Ala Val Asp Leu Val Ser  
340 345 350

Arg Leu Leu Gln Tyr Ser Pro Asn Leu Arg Cys Thr Val Leu Asp Ala  
355 360 365

Leu Asp Ala Pro Phe Pro Leu Asp Glu Phe Arg Asp Pro Asn Pro Arg  
370 375 380

Leu Pro Asn Gly Pro Ile Xaa Pro Thr Thr Ile Asn Ser Asn Pro  
385 390 395

<210> 15

<211> 1673

<212> DNA

<213> Triticum aestivum

<220>

<221> unsure

<222> (1349)

<400> 15

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ctccaccacc atcggcggca agaacggcga gcccaagcag acgattagct acatggcgga 240  
gcgcgttgg 99actgggtt cgtttggcat cgtcttttag gctaaatgcc tggaaaccgg 300  
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gcttatgcgt tcgatgatcc attccaatgt tgcgtccctc aagcactgct tcttctcaac 420  
cacaagtaga gatgagatgt tcctgaacct tgcgtcatggag tatgtcccg agacgctata 480  
cccgctgctt aagcaactaca gtaatgccaa ccagggatg ccgccttatct atgtcaagct 540  
ttacatgtat cagctttta gagggcttagc ttatgttcat actgttccag gagtttgcca 600  
cagggatgtg aaaccacaaa atgttttgt tgatcctcta acccatcaag tcaagatctg 660  
tgactttgg aatgcacaaat ttctggtacc tggtaaccc aacatagcat acatatgctc 720  
tcgctactat cgtgctcctg agctcatatt tggtaact gaatatacaa cttcaataga 780  
catatggtca gctggatgtt ttcttgccaga gctacttctt ggtcagccctc tgtttccagg 840  
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tccttggcac aagattttcc acaagagaat gcccgtgaa gctatagatc ttgcctcccg 1020  
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ggAACATGTT CGACGGCAAA ATGGCCCAA CTTCGCCCAT GCTGGGAGCT AAACGGGGCG 1260  
CGCCCGCATTGCCCATATT TTGTTGTCC GCCATCATCG AAGAATCAAT CTCTCCCTA 1320  
AATCCTGAGG AGAGACCGAT CAAGTGCANT GCCAGTGCCA GTGAAAGAAG TACAACTATG 1380  
TAAATTACCT GACCTTGGAA GAATCGTTGT TGTTGTTGCC GGTGCCGGCC ATGTTAAGT 1440  
ACATGGCGC ACATGTTGGT TGAGTGTAA CCTATTATTA AGTAGGTAAG AGCAATGATG 1500  
TAGGAGGTGG AGACATATGT TAATGCTAGG TCTGTGACCT GTTTAAGTA CATTTTGTA 1560  
ATGCTTGGTA GTGGTACTGT AATGCGGCAA TAGCTGCTCC ATGTTTGTG CTTGTCCT 1620  
GATGAAATG TCGTCGTCCT GCAGCAAAAAA AAAAAAAAAAAA AAAA 1673

*Groh*  
*B13*

<210> 16  
<211> 402  
<212> PRT  
<213> Triticum aestivum

<400> 16  
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Pro Pro Thr Ala Val Ala Cys Glu Lys Lys Gln Gln Asp Gly Glu Ala  
20 25 30  
Pro Tyr Ala Glu Gly Asn Asp Ala Met Thr Gly His Ile Ile Ser Thr  
35 40 45  
Thr Ile Gly Gly Lys Asn Gly Glu Pro Lys Gln Thr Ile Ser Tyr Met  
50 55 60  
Ala Glu Arg Val Val Gly Thr Gly Ser Phe Gly Ile Val Phe Gln Ala  
65 70 75 80  
Lys Cys Leu Glu Thr Gly Glu Met Val Gly Ile Lys Lys Val Leu Gln  
85 90 95  
Asp Arg Arg Tyr Lys Asn Arg Glu Leu Gln Leu Met Arg Ser Met Ile  
100 105 110  
His Ser Asn Val Val Ser Leu Lys His Cys Phe Phe Ser Thr Thr Ser  
115 120 125  
Arg Asp Glu Leu Phe Leu Asn Leu Val Met Glu Tyr Val Pro Glu Thr  
130 135 140  
Leu Tyr Arg Val Leu Lys His Tyr Ser Asn Ala Asn Gln Gly Met Pro  
145 150 155 160  
Leu Ile Tyr Val Lys Leu Tyr Met Tyr Gln Leu Phe Arg Gly Leu Ala  
165 170 175  
Tyr Val His Thr Val Pro Gly Val Cys His Arg Asp Val Lys Pro Gln  
180 185 190  
Asn Val Leu Val Asp Pro Leu Thr His Gln Val Lys Ile Cys Asp Phe  
195 200 205  
Gly Ser Ala Lys Val Leu Val Pro Gly Glu Pro Asn Ile Ala Tyr Ile  
210 215 220  
Cys Ser Arg Tyr Tyr Arg Ala Pro Glu Leu Ile Phe Gly Ala Thr Glu  
225 230 235 240  
Tyr Thr Thr Ser Ile Asp Ile Trp Ser Ala Gly Cys Val Leu Ala Glu  
245 250 255  
Leu Leu Leu Gly Gln Pro Leu Phe Pro Gly Glu Thr Ala Val Asp Gln  
260 265 270  
Leu Val Glu Ile Ile Lys Val Leu Gly Thr Pro Thr Arg Glu Glu Ile  
275 280 285

Arg Cys Met Asn Pro Asn Tyr Thr Glu Phe Arg Phe Pro Gln Ile Lys  
290 295 300

Ala His Pro Trp His Lys Ile Phe His Lys Arg Met Pro Ala Glu Ala  
305 310 315 320

Ile Asp Leu Ala Ser Arg Leu Leu Gln Tyr Ser Pro Asn Leu Arg Cys  
325 330 335

Thr Ala Leu Asp Ala Cys Ala His Ser Phe Phe Asp Glu Leu Arg Glu  
340 345 350

Pro Asn Ala Arg Leu Pro Asn Gly Arg Pro Phe Pro Pro Leu Phe Asn  
355 360 365

Phe Lys Pro Glu Leu Ala Asn Ala Ser Pro Glu Leu Ile Asn Arg Leu  
370 375 380

Val Pro Glu His Val Arg Arg Gln Asn Gly Pro Asn Phe Ala His Ala  
385 390 395 400

Gly Ser

<210> 17

<211> 639

<212> PRT

<213> Zea mays

<400> 17

Met Gly Asn Thr Cys Val Gly Pro Ser Ile Thr Met Asn Gly Phe Phe  
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Gln Ser Val Ser Thr Ala Leu Trp Lys Thr Pro Gln Glu Gly Asp Ala  
20 25 30

Leu Pro Ala Ala Ala Asn Gly Pro Gly Gly Pro Ala Gly Ala Gly Ser  
35 40 45

Gln Ser Ala Leu Pro Lys Pro Ala Ser Asp Val His His Val Ala Val  
50 55 60

Gln Ser Glu Ala Pro Glu Pro Val Lys Ile Ala Ala Tyr His Ser Glu  
65 70 75 80

Pro Ala Pro Ala Val Arg Ser Glu Ala Pro Glu Pro Val Lys Ile Ala  
85 90 95

Ala Ser His Ser Glu Pro Ala Pro Met Ala Ala Lys Pro Gly Gly Ala  
100 105 110

Ala Ala Asn Ala Ser Pro Ser Pro Ser Pro Arg Pro Arg Pro Gln Val  
115 120 125

Lys Arg Val Ser Ser Ala Gly Leu Leu Leu Gly Ser Val Leu Arg Arg  
130 135 140

Lys Thr Glu Asn Leu Lys Asp Lys Tyr Ser Leu Gly Arg Arg Leu Gly  
145 150 155 160

Gln Gly Gln Phe Gly Thr Thr His Leu Cys Val Glu Arg Ala Thr Gly  
165 170 175

Lys Glu Leu Ala Cys Lys Ser Ile Leu Lys Arg Lys Leu Gly Ser Asp  
180 185 190

Asp Asp Val Glu Asp Val Arg Arg Glu Ile Gln Ile Met His His Leu  
195 200 205

Ala Gly His Pro Ser Val Val Gly Ile Arg Gly Ala Tyr Glu Asp Ala  
210 215 220

Val Ala Val His Leu Val Met Glu Leu Cys Gly Gly Gly Glu Leu Phe  
225 230 235 240

Asp Arg Ile Val Arg Arg Gly His Tyr Thr Glu Arg Lys Ala Ala Glu  
245 250 255

Leu Ala Arg Val Ile Val Gly Val Val Glu Ala Cys His Ser Met Gly  
260 265 270

Val Met His Arg Asp Leu Lys Pro Glu Asn Phe Leu Phe Ala Asp His  
275 280 285

Ser Glu Glu Ala Ala Leu Lys Thr Ile Asp Phe Gly Leu Ser Ile Phe  
290 295 300

Phe Arg Pro Gly Gln Ile Phe Thr Asp Val Val Gly Ser Pro Tyr Tyr  
305 310 315 320

Val Ala Pro Glu Val Leu Lys Lys Arg Tyr Gly Pro Glu Ala Asp Val  
325 330 335

Trp Ser Ala Gly Val Ile Ile Tyr Ile Leu Leu Cys Gly Val Pro Pro  
340 345 350

Phe Trp Ala Glu Asn Glu Gln Gly Ile Phe Glu Glu Val Leu His Gly  
355 360 365

Arg Leu Asp Phe Glu Ser Glu Pro Trp Pro Ser Ile Ser Asp Gly Ala  
370 375 380

Lys Asp Leu Val Arg Arg Met Leu Val Arg Asp Pro Arg Lys Arg Leu  
385 390 395 400

Thr Ala His Glu Val Leu Arg His Pro Trp Val Gln Val Gly Gly Val  
405 410 415

Ala Pro Asp Arg Pro Leu Asp Ser Ala Val Leu Ser Arg Met Lys Gln  
420 425 430

Phe Ser Ala Met Asn Lys Leu Lys Lys Met Ala Leu Arg Val Ile Ala  
435 440 445

Glu Asn Leu Ser Glu Asp Glu Ile Ala Gly Leu Arg Glu Met Phe Lys  
450 455 460

Met Ile Asp Ala Asp Asn Ser Gly Gln Ile Thr Phe Glu Glu Leu Lys  
465 470 475 480

Val Gly Leu Glu Lys Val Gly Ala Asn Leu Gln Glu Ser Glu Ile Tyr  
485 490 495

Ala Leu Met Gln Ala Ala Asp Val Asp Asn Asn Gly Thr Ile Asp Tyr  
500 505 510

Gly Glu Phe Ile Ala Ala Thr Leu His Leu Asn Lys Val Glu Arg Glu  
515 520 525

Asp His Leu Phe Ala Ala Phe Gln Tyr Phe Asp Lys Asp Gly Ser Gly  
530 535 540

Tyr Ile Thr Ala Asp Glu Leu Gln Val Ala Cys Glu Glu Phe Gly Leu  
545 550 555 560

Gly Asp Val Gln Leu Glu Asp Leu Ile Gly Glu Val Asp Gln Asp Asn  
565 570 575

Asp Gly Arg Ile Asp Tyr Asn Glu Phe Val Ala Met Met Gln Lys Pro  
580 585 590

Thr Val Gly Gly Ser Arg Arg Arg Pro Ile Cys Arg Thr Ala Ser Ala  
595 600 605

Ser Gly Ser Ala Ser Gly Ser Gly Arg Arg Ser Gly Trp Pro Arg Pro  
610 615 620

Leu Cys Leu Trp Leu Pro Cys Cys Leu Arg Val Gly Val Asp Asp  
625 630 635

<210> 18  
<211> 625  
<212> PRT  
<213> Zea mays

<400> 18  
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His Asp Ala Asp Pro Ser Gly Ala Gly Ser Val Ala Pro Pro Ser Pro  
20 25 30

Leu Pro Ala Asp Gly Ala Pro Leu Pro Ala Thr Pro Arg Arg His Lys  
35 40 45

Ser Gly Ser Thr Thr Pro Val His His His Gln Ala Ala Thr Pro Gly  
50 55 60

Ala Ala Ala Trp Pro Ser Pro Tyr Pro Ala Gly Gly Ala Ser Pro Leu  
65 70 75 80

Pro Ala Gly Val Ser Pro Ser Pro Ala Arg Ser Thr Pro Arg Arg Phe  
85 90 95

Phe Lys Arg Pro Phe Pro Pro Ser Pro Ala Lys His Ile Lys Ala  
100 105 110

Thr Leu Ala Lys Arg Leu Gly Gly Lys Pro Lys Glu Gly Thr Ile  
115 120 125

*Cat  
B*

Pro Glu Glu Gly Gly Ala Gly Ala Gly Ala Gly Ala Gly Ala  
130 135 140

Gly Ala Ala Val Gly Ala Ala Asp Ser Ala Glu Ala Asp Arg Pro Leu  
145 150 155 160

Asp Lys Thr Phe Gly Phe Ala Lys Asn Phe Gly Ala Lys Tyr Asp Leu  
165 170 175

Gly Lys Glu Val Gly Arg Gly His Phe Gly His Thr Cys Ser Ala Val  
180 185 190

Val Lys Lys Gly Glu His Lys Gly His Thr Val Ala Val Lys Ile Ile  
195 200 205

Ser Lys Ala Lys Met Thr Thr Ala Ile Ser Ile Glu Asp Val Arg Arg  
210 215 220

Glu Val Lys Ile Leu Lys Ala Leu Ser Gly His Asp Asp Leu Val Arg  
225 230 235 240

Phe Tyr Asp Ala Cys Glu Asp Ala Leu Asn Val Tyr Ile Val Met Glu  
245 250 255

Leu Cys Glu Gly Glu Leu Leu Asp Arg Ile Leu Ala Arg Gly Gly  
260 265 270

Arg Tyr Thr Glu Glu Asp Ala Lys Ala Ile Ile Val Gln Ile Leu Ser  
275 280 285

Val Val Ala Phe Cys His Leu Gln Gly Val Val His Arg Asp Leu Lys  
290 295 300

Pro Glu Asn Phe Leu Phe Thr Thr Arg Asp Glu Ser Ala Pro Met Lys  
305 310 315 320

Leu Ile Asp Phe Gly Leu Ser Asp Phe Ile Arg Pro Asp Glu Arg Leu  
325 330 335

Asn Asp Ile Val Gly Ser Ala Tyr Tyr Val Ala Pro Glu Val Leu His  
340 345 350

Arg Ser Tyr Ser Met Glu Ala Asp Ile Trp Ser Ile Gly Val Ile Thr  
355 360 365

Tyr Ile Leu Leu Cys Gly Ser Arg Pro Phe Trp Ala Arg Thr Glu Ser  
370 375 380

Gly Ile Phe Arg Ser Val Leu Arg Ala Asp Pro Asn Phe Asp Asp Ser  
385 390 395 400

Pro Trp Pro Ser Val Ser Ala Glu Ala Lys Asp Phe Val Lys Arg Phe  
405 410 415

Leu Asn Lys Asp Tyr Arg Lys Arg Met Thr Ala Val Gln Ala Leu Thr  
420 425 430

His Pro Trp Leu Arg Asp Glu Gln Arg Gln Ile Pro Leu Asp Ile Leu  
435 440 445

DOCUMENTATION

*BBB*

Ile Phe Arg Leu Val Lys Gln Tyr Leu Arg Ala Thr Pro Leu Lys Arg  
450 455 460

Leu Ala Leu Lys Ala Leu Ser Lys Ala Leu Ser Glu Asp Glu Leu Leu  
465 470 475 480

Tyr Leu Arg Leu Gln Phe Lys Leu Leu Glu Pro Arg Asp Gly Phe Val  
485 490 495

Ser Leu Asp Asn Phe Arg Thr Ala Leu Thr Arg Tyr Ser Thr Asp Ala  
500 505 510

Met Arg Glu Ser Arg Val Leu Glu Phe Gln His Ala Leu Glu Pro Leu  
515 520 525

Ala Tyr Arg Lys Met Asp Phe Glu Glu Phe Cys Ala Ala Ala Ile Ser  
530 535 540

Pro Tyr Gln Leu Glu Ala Leu Glu Arg Trp Glu Glu Ile Ala Gly Thr  
545 550 555 560

Ala Phe Gln His Phe Glu Gln Glu Gly Asn Arg Val Ile Ser Val Glu  
565 570 575

Glu Leu Ala Gln Glu Leu Asn Leu Ala Pro Thr His Tyr Ser Ile Val  
580 585 590

Gln Asp Trp Ile Arg Lys Ser Asp Gly Lys Leu Asn Phe Leu Gly Phe  
595 600 605

Thr Lys Phe Leu His Gly Val Thr Ile Arg Gly Ser Asn Thr Arg Arg  
610 615 620

His  
625

<210> 19  
<211> 576  
<212> PRT  
<213> *Arabidopsis thaliana*

<400> 19  
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Pro Val Ser Gly Glu Thr Asn Glu Ala Pro Thr Asn Ser Gln Pro Pro  
20 25 30

Ala Lys Ser Ser Gly Phe Pro Phe Tyr Ser Pro Ser Pro Val Pro Ser  
35 40 45

Leu Phe Lys Ser Ser Pro Ser Val Ser Ser Ser Val Ser Ser Thr Pro  
50 55 60

Leu Arg Ile Phe Lys Arg Pro Phe Pro Pro Ser Pro Ala Lys His  
65 70 75 80

Ile Arg Ala Phe Leu Ala Arg Arg Tyr Gly Ser Val Lys Pro Asn Glu  
85 90 95

Val Ser Ile Pro Glu Gly Lys Glu Cys Glu Ile Gly Leu Asp Lys Ser  
 100 105 110

Phe Gly Phe Ser Lys Gln Phe Ala Ser His Tyr Glu Ile Asp Gly Glu  
 115 120 125

*Cat 3/13*  
 Val Gly Arg Gly His Phe Gly Tyr Thr Cys Ser Ala Lys Gly Lys Lys  
 130 135 140

Gly Ser Leu Lys Gly Gln Glu Val Ala Val Lys Val Ile Pro Lys Ser  
 145 150 155 160

Lys Met Thr Thr Ala Ile Ala Ile Glu Asp Val Ser Arg Glu Val Lys  
 165 170 175

Met Leu Arg Ala Leu Thr Gly His Lys Asn Leu Val Gln Phe Tyr Asp  
 180 185 190

Ala Phe Glu Asp Asp Glu Asn Val Tyr Ile Val Met Glu Leu Cys Lys  
 195 200 205

Gly Gly Glu Leu Leu Asp Lys Ile Leu Gln Arg Gly Gly Lys Tyr Ser  
 210 215 220

Glu Asp Asp Ala Lys Lys Val Met Val Gln Ile Leu Ser Val Val Ala  
 225 230 235 240

Tyr Cys His Leu Gln Gly Val Val His Arg Asp Leu Lys Pro Glu Asn  
 245 250 255

Phe Leu Phe Ser Thr Lys Asp Glu Thr Ser Pro Leu Lys Ala Ile Asp  
 260 265 270

Phe Gly Leu Ser Asp Tyr Val Lys Pro Asp Glu Arg Leu Asn Asp Ile  
 275 280 285

Val Gly Ser Ala Tyr Tyr Val Ala Pro Glu Val Leu His Arg Thr Tyr  
 290 295 300

Gly Thr Glu Ala Asp Met Trp Ser Ile Gly Val Ile Ala Tyr Ile Leu  
 305 310 315 320

Leu Cys Gly Ser Arg Pro Phe Trp Ala Arg Thr Glu Ser Gly Ile Phe  
 325 330 335

Arg Ala Val Leu Lys Ala Glu Pro Asn Phe Glu Glu Ala Pro Trp Pro  
 340 345 350

Ser Leu Ser Pro Glu Ala Val Asp Phe Val Lys Arg Leu Leu Asn Lys  
 355 360 365

Asp Tyr Arg Lys Arg Leu Thr Ala Ala Gln Ala Leu Cys His Pro Trp  
 370 375 380

Leu Val Gly Ser His Glu Leu Lys Ile Pro Ser Asp Met Ile Ile Tyr  
 385 390 395 400

Lys Leu Val Lys Val Tyr Ile Met Ser Thr Ser Leu Arg Lys Ser Ala  
 405 410 415

Leu Ala Ala Leu Ala Lys Thr Leu Thr Val Pro Gin Leu Ala Tyr Leu  
420 425 430

Arg Glu Gln Phe Thr Leu Leu Gly Pro Ser Lys Asn Gly Tyr Ile Ser  
435 440 445

*Gut*  
*B13*

Met Gln Asn Tyr Lys Thr Ala Ile Leu Lys Ser Ser Thr Asp Ala Met  
450 455 460

Lys Asp Ser Arg Val Phe Asp Phe Val His Met Ile Ser Cys Leu Gln  
465 470 475 480

Tyr Lys Lys Leu Asp Phe Glu Glu Phe Cys Ala Ser Ala Leu Ser Val  
485 490 495

Tyr Gln Leu Glu Ala Met Glu Thr Trp Glu Gln His Ala Arg Arg Ala  
500 505 510

Tyr Glu Leu Phe Glu Lys Asp Gly Asn Arg Pro Ile Met Ile Glu Glu  
515 520 525

Leu Ala Ser Glu Leu Gly Leu Gly Pro Ser Val Pro Val His Val Val  
530 535 540

Leu Gln Asp Trp Ile Arg His Ser Asp Gly Lys Leu Ser Phe Leu Gly  
545 550 555 560

Phe Val Arg Leu Leu His Gly Val Ser Ser Arg Thr Leu Gln Lys Ala  
565 570 575

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<211> 405  
<212> PRT  
<213> Arabidopsis thaliana

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Gly Val Asp Lys Leu Pro Glu Glu Met Asn Asp Met Lys Ile Arg Asp  
20 25 30

Asp Lys Glu Met Glu Ala Thr Val Val Asp Gly Asn Gly Thr Glu Thr  
35 40 45

Gly His Ile Ile Val Thr Thr Ile Gly Gly Arg Asn Gly Gln Pro Lys  
50 55 60

Gln Thr Ile Ser Tyr Met Ala Glu Arg Val Val Gly His Gly Ser Phe  
65 70 75 80

Gly Val Val Phe Gln Ala Lys Cys Leu Glu Thr Gly Glu Thr Val Ala  
85 90 95

Ile Lys Lys Val Leu Gln Asp Arg Arg Tyr Lys Asn Arg Glu Leu Gln  
100 105 110

Thr Met Arg Leu Leu Asp His Pro Asn Val Val Ser Leu Lys His Cys  
115 120 125

DRAFT FEB 2000

Phe Phe Ser Thr Thr Glu Lys Asp Glu Leu Tyr Leu Asn Leu Val Leu  
130 135 140

Glu Tyr Val Pro Glu Thr Val His Arg Val Ile Lys His Tyr Asn Lys  
145 150 155 160

*Cat*  
*B13*

Leu Asn Gln Arg Met Pro Leu Ile Tyr Val Lys Leu Tyr Thr Tyr Gln  
165 170 175

Ile Phe Arg Ala Leu Ser Tyr Ile His Arg Cys Ile Gly Val Cys His  
180 185 190

Arg Asp Ile Lys Pro Gln Asn Leu Leu Val Asn Pro His Thr His Gln  
195 200 205

Val Lys Leu Cys Asp Phe Gly Ser Ala Lys Val Leu Val Lys Gly Glu  
210 215 220

Pro Asn Ile Ser Tyr Ile Cys Ser Arg Tyr Tyr Arg Ala Pro Glu Leu  
225 230 235 240

Ile Phe Gly Ala Thr Glu Tyr Thr Thr Ala Ile Asp Val Trp Ser Ala  
245 250 255

Gly Cys Val Leu Ala Glu Leu Leu Gly Gln Pro Leu Phe Pro Gly  
260 265 270

Glu Ser Gly Val Asp Gln Leu Val His Ile Ile Lys Val Leu Gly Thr  
275 280 285

Pro Thr Arg Glu Glu Ile Lys Cys Met Asn Pro Asn Tyr Thr Glu Phe  
290 295 300

Lys Phe Pro Gln Ile Lys Ala His Pro Trp His Lys Ile Phe His Lys  
305 310 315 320

Arg Met Pro Pro Glu Ala Val Asp Leu Val Ser Arg Leu Leu Gln Tyr  
325 330 335

Ser Pro Asn Leu Arg Ser Ala Ala Leu Asp Thr Leu Val His Pro Phe  
340 345 350

Phe Asp Glu Leu Arg Asp Pro Asn Ala Arg Leu Pro Asn Gly Arg Phe  
355 360 365

Leu Pro Pro Ala Phe His Phe Lys Pro His Glu Leu Lys Gly Val Pro  
370 375 380

Leu Glu Met Val Ala Lys Leu Val Pro Glu His Ala Arg Lys Gln Cys  
385 390 395 400

Pro Trp Leu Gly Leu  
405

<210> 21  
<211> 412  
<212> PRT  
<213> *Medicago sativa*

TOP F20-T215800

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Asn Ala Ser Ser Val Gly Val Glu Lys Leu Pro Glu Glu Met Asn Asp  
20 25 30  
  
Met Lys Ile Arg Asp Asp Lys Glu Met Glu Ala Ala Thr Ile Val Asp  
35 40 45  
  
Gly Asn Gly Thr Glu Thr Gly His Ile Ile Val Thr Thr Ile Gly Gly  
50 55 60  
  
Lys Asn Gly Gln Pro Lys Gln Thr Ile Ser Tyr Met Ala Glu Arg Val  
65 70 75 80  
  
Val Gly His Gly Ser Phe Gly Val Val Phe Gln Ala Lys Cys Leu Glu  
85 90 95  
  
Thr Gly Glu Thr Val Ala Ile Lys Lys Val Leu Gln Asp Lys Arg Tyr  
100 105 110  
  
Lys Asn Arg Glu Leu Gln Thr Met Arg Leu Leu Asp His Pro Asn Val  
115 120 125  
  
Val Ser Leu Lys His Cys Phe Phe Ser Thr Thr Glu Lys Asp Glu Leu  
130 135 140  
  
Tyr Leu Asn Leu Val Leu Glu Tyr Val Pro Glu Thr Val Ser Arg Val  
145 150 155 160  
  
Ile Arg His Tyr Asn Lys Met Asn Gln Arg Met Pro Met Ile Tyr Val  
165 170 175  
  
Lys Leu Tyr Ser Tyr Gln Ile Cys Arg Ala Leu Ala Tyr Ile His Asn  
180 185 190  
  
Ser Ile Gly Val Cys His Arg Asp Ile Lys Pro Gln Asn Leu Leu Val  
195 200 205  
  
Asn Pro His Thr His Gln Leu Lys Ile Cys Asp Phe Gly Ser Ala Lys  
210 215 220  
  
Val Leu Val Lys Gly Glu Pro Asn Ile Ser Tyr Ile Cys Ser Arg Tyr  
225 230 235 240  
  
Tyr Arg Ala Pro Glu Leu Ile Phe Gly Ala Thr Glu Tyr Thr Thr Ala  
245 250 255  
  
Ile Asp Ile Trp Ser Ala Gly Cys Val Leu Gly Glu Leu Leu Gly  
260 265 270  
  
Gln Pro Leu Phe Pro Gly Glu Ser Gly Val Asp Gln Leu Val Glu Ile  
275 280 285  
  
Ile Lys Val Leu Gly Thr Pro Thr Arg Glu Glu Ile Lys Cys Met Asn  
290 295 300  
  
Pro Asn Tyr Thr Glu Phe Lys Phe Pro Gln Ile Lys Ala His Pro Trp  
305 310 315 320

His Lys Ile Phe His Lys Arg Met Pro Pro Glu Ala Val Asp Leu Val  
325 330 335

Ser Arg Leu Leu Gln Tyr Ser Pro Asn Leu Arg Ser Thr Ala Leu Glu  
340 345 350

Ala Leu Val His Pro Phe Tyr Asp Asp Val Arg Asp Pro Asn Thr Arg  
355 360 365

Leu Pro Asn Gly Arg Phe Leu Pro Pro Leu Phe Asn Phe Lys Val Asn  
370 375 380

Glu Leu Lys Gly Val Pro Ala Glu Met Leu Val Lys Leu Val Pro Pro  
385 390 395 400

His Ala Arg Lys Gln Cys Ala Leu Phe Gly Ser Ser  
405 410

<210> 22

<211> 411

<212> PRT

<213> Medicago sativa

<400> 22

Met Ala Ser Val Gly Val Ala Pro Thr Ser Gly Phe Arg Glu Val Leu  
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Gly Asp Gly Glu Ile Gly Val Asp Asp Ile Leu Pro Glu Glu Met Ser  
20 25 30

Asp Met Lys Ile Arg Asp Asp Arg Glu Met Glu Ala Thr Val Val Asp  
35 40 45

Gly Asn Gly Thr Glu Thr Gly His Ile Ile Val Thr Thr Ile Gly Gly  
50 55 60

Arg Asn Gly Gln Pro Lys Gln Thr Ile Ser Tyr Met Ala Glu Arg Val  
65 70 75 80

Val Gly His Gly Ser Phe Gly Val Val Phe Gln Ala Lys Cys Leu Glu  
85 90 95

Thr Gly Glu Thr Val Ala Ile Lys Lys Val Leu Gln Asp Lys Arg Tyr  
100 105 110

Lys Asn Arg Glu Leu Gln Thr Met Arg Leu Leu Asp His Pro Asn Val  
115 120 125

Val Ser Leu Lys His Cys Phe Phe Ser Thr Thr Glu Lys Asp Glu Leu  
130 135 140

Tyr Leu Asn Leu Val Leu Glu Tyr Val Pro Glu Thr Val His Arg Val  
145 150 155 160

Ile Lys His Tyr Ser Lys Leu Asn Gln Arg Met Pro Met Ile Tyr Val  
165 170 175

Lys Leu Tyr Thr Tyr Gln Ile Phe Arg Ala Leu Ser Tyr Ile His Arg  
180 185 190

Cys Ile Gly Val Cys His Arg Asp Ile Lys Pro Gln Asn Leu Leu Val  
195 200 205

Asn Pro His Thr His Gln Val Lys Leu Cys Asp Phe Gly Ser Ala Lys  
210 215 220

Val Leu Val Lys Gly Glu Pro Asn Ile Ser Tyr Ile Cys Ser Arg Tyr  
225 230 235 240

Tyr Arg Ala Pro Glu Leu Ile Phe Gly Ala Thr Glu Tyr Thr Thr Ala  
245 250 255

Ile Asp Val Trp Ser Val Gly Cys Val Leu Ala Glu Leu Leu Leu Gly  
260 265 270

Gln Pro Leu Phe Pro Gly Glu Arg Gly Val Asp Gln Leu Val Glu Ile  
275 280 285

Ile Lys Val Leu Gly Thr Pro Thr Arg Glu Glu Ile Lys Cys Met Asn  
290 295 300

Pro Asn Tyr Thr Glu Phe Lys Phe Pro Gln Ile Lys Ala His Pro Trp  
305 310 315 320

His Lys Ile Phe His Lys Arg Met Pro Ala Glu Ala Val Asp Leu Val  
325 330 335

Ser Arg Leu Leu Gln Tyr Ser Pro Asn Leu Arg Cys Gln Ala Leu Asp  
340 345 350

Cys Leu Thr His Pro Phe Phe Asp Glu Leu Arg Asp Pro Asn Ala Arg  
355 360 365

Leu Pro Thr Gly Arg Phe Leu Pro Pro Leu Phe Asn Phe Lys Pro His  
370 375 380

Glu Leu Lys Gly Val Pro Val Glu Thr Leu Met Lys Leu Val Pro Glu  
385 390 395 400

His Ala Arg Lys Gln Cys Pro Phe Leu Gly Leu  
405 410

<210> 23

<211> 407

<212> PRT

<213> *Arabidopsis thaliana*

<400> 23

Met Ala Ser Leu Pro Leu Gly Pro Gln Pro His Ala Leu Ala Pro Pro  
1 5 10 15

Leu Gln Leu His Asp Gly Asp Ala Leu Lys Arg Arg Pro Glu Leu Asp  
20 25 30

Ser Asp Lys Glu Met Ser Ala Ala Val Ile Glu Gly Asn Asp Ala Val  
35 40 45

Thr Gly His Ile Ile Ser Thr Thr Ile Gly Gly Lys Asn Gly Glu Pro  
50 55 60

Lys Gln Thr Ile Ser Tyr Met Ala Glu Arg Val Val Gly Thr Gly Ser  
65 70 75 80

Phe Gly Ile Val Phe Gln Ala Lys Cys Leu Glu Thr Gly Glu Ser Val  
85 90 95

Ala Ile Lys Lys Val Leu Gln Asp Arg Arg Tyr Lys Asn Arg Glu Leu  
100 105 110

Gln Leu Met Arg Pro Met Asp His Pro Asn Val Ile Ser Leu Lys His  
115 120 125

Cys Phe Phe Ser Thr Thr Ser Arg Asp Glu Leu Phe Leu Asn Leu Val  
130 135 140

Met Glu Tyr Val Pro Glu Thr Leu Tyr Arg Val Leu Arg His Tyr Thr  
145 150 155 160

Ser Ser Asn Gln Arg Met Pro Ile Phe Tyr Val Lys Leu Tyr Thr Tyr  
165 170 175

Gln Ile Phe Arg Gly Leu Ala Tyr Ile His Thr Val Pro Gly Val Cys  
180 185 190

His Arg Asp Val Lys Pro Gln Asn Leu Leu Val Asp Pro Leu Thr His  
195 200 205

Gln Val Lys Leu Cys Asp Phe Gly Ser Ala Lys Val Leu Val Lys Gly  
210 215 220

Glu Pro Asn Ile Ser Tyr Ile Cys Ser Arg Tyr Tyr Arg Ala Pro Glu  
225 230 235 240

Leu Ile Phe Gly Ala Thr Glu Tyr Thr Ala Ser Ile Asp Ile Trp Ser  
245 250 255

Ala Gly Cys Val Leu Ala Glu Leu Leu Leu Gly Gln Pro Leu Phe Pro  
260 265 270

Gly Glu Asn Ser Val Asp Gln Leu Val Glu Ile Ile Lys Val Leu Gly  
275 280 285

Thr Pro Thr Arg Glu Glu Ile Arg Cys Met Asn Pro Asn Tyr Thr Asp  
290 295 300

Phe Arg Phe Pro Gln Ile Lys Ala His Pro Trp His Lys Val Phe His  
305 310 315 320

Lys Arg Met Pro Pro Glu Ala Ile Asp Leu Ala Ser Arg Leu Leu Gln  
325 330 335

Tyr Ser Pro Ser Leu Arg Cys Thr Ala Leu Glu Ala Cys Ala His Pro  
340 345 350

Phe Phe Asn Glu Leu Arg Glu Pro Asn Ala Arg Leu Pro Asn Gly Arg  
355 360 365

Pro Leu Pro Pro Leu Phe Asn Phe Lys Gln Glu Leu Gly Gly Ala Ser  
370 375 380

Met Glu Leu Ile Asn Arg Leu Ile Pro Glu His Val Arg Arg Gln Met  
385 390 395 400

Ser Thr Gly Leu Gln Asn Ser  
405

Cont  
B13

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